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DATE: Wednesday, September 14, 2005

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	DB=PG	PB, USPT; PLUR=YES; OP=ADJ	
	L6	tsi-a or tsi-b	1
	L5	transcriptionally silent information	0
	L4	L3 and impair\$	161
	L3	L2 and select\$ [clm]	415
	L2	L1 and plant	595
	L1	transcriptional gene silenc\$	658

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=> s transcriptional gene silenc? 1038 TRANSCRIPTIONAL GENE SILENC?

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L2
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760 L1 AND PLANT?

- => del 13. DELETE L3? (Y)/N:y
- => s l2 and (defect? or impair? or muta?)
 L3 168 L2 AND (DEFECT? OR IMPAIR? OR MUTA?)
- => s 13 and select?

L4 17 L3 AND SELECT?

=> dup rem 14 PROCESSING COMPLETED FOR L4

L5 10 DUP REM L4 (7 DUPLICATES REMOVED)

=> d 1-10 ti

- L5 ANSWER 1 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1
- TI UPS1 and UPS2 from Arabidopsis Mediate High Affinity Transport of Uracil and 5-Fluorouracil
- L5 ANSWER 2 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 2
- TI Inhibition of HIV-1 fusion with small interfering RNAs targeting the chemokine coreceptor CXCR4
- L5 ANSWER 3 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3
- TI Tandem inverted repeat system for **selection** of effective transgenic RNAi strains in chlamydomonas
- L5 ANSWER 4 OF 10 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN DUPLICATE 4
- TI Short defective interfering RNAs of tombusviruses are not targeted but trigger post-transcriptional gene silencing against their helper virus.
- L5 ANSWER 5 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI High-oleic and high-stearic cottonseed oils: nutritionally improved cooking oils developed using gene silencing
- L5 ANSWER 6 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5
- TI Virus variation in relation to resistance-breaking in plants
- L5 ANSWER 7 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI The abundant retinal protein of the Chlamydomonas eye is not the photoreceptor for phototaxis and photophobic responses.
- L5 ANSWER 8 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI A library of Arabidopsis 35S-cDNA lines for identifying novel mutants
- L5 ANSWER 9 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Selective inhibition of gene expression by RNAi in chick embryos in ovo
- L5 ANSWER 10 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- TI Cytosine methylation at CG and CNG sites is not a prerequisite for the initiation of transcriptional gene silencing in plants, but it is required for its maintenance
- => d ab
- L5 ANSWER 1 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1
- AB Salvage pathways play an important role in providing nucleobases to cells,

which are unable to synthesize sufficient amts. for their needs. Cellular uptake systems for pyrimidines have been described, but in higher eukaryotes, transporters for thymine and uracil have not been identified. Two plant transporters, AtUPS1 and PvUPS1, were recently identified as transporters for allantoin in Arabidopsis and French bean, resp. However, Arabidopsis, in contrast to tropical legumes, uses mainly amino acids for long distance transport. Allantoin transport has not been described in the Brassicaceae. Thus, the physiol. substrates of ureide permease (UPS) transporters in Arabidopsis may be compds. structurally related to allantoin. A detailed anal. of the substrate specificities of two members of the AtUPS family shows that AtUPS1 and AtUPS2 mediate high affinity uracil and 5-fluorouracil (a toxic uracil analog) transport when expressed in yeast and Xenopus oocytes. Consistent with a function during germination and early seedling development, AtUPS1 expression is transiently induced during the early stages of seedling development followed by up-regulation of AtUPS2 expression. Arabidopsis ups2 insertion mutants and ups1 lines, in which transcript levels were reduced by post-transcriptional gene silencing, are more tolerant to 5-fluorouracil as compared with wild type plants. The results suggest that in Arabidopsis UPS transporters are the main transporters for uracil and potentially other nucleobases, whereas during evolution legumes may have taken advantage of the low selectivity of UPS proteins for long distance transport of allantoin.

=> d so

- L5 ANSWER 1 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1 SO Journal of Biological Chemistry (2004), 279(43), 44817-44824 CODEN: JBCHA3; ISSN: 0021-9258
- => d 3 ab
- L5 ANSWER 3 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3 RNA interference (RNAi), the double-stranded RNA (dsRNA) triggered post-AΒ transcriptional gene silencing, is becoming a powerful tool for reverse genetics studies. Stable RNAi, induced by the expression of inverted repeat (IR) transgenes, has been achieved in protozoa, algae, fungi, plants, and metazoans. However, the level of gene silencing is often quite variable, depending on the type of construct, transgene copy number, site of integration, and target gene. is a hindrance in functional genomics studies, where it is desirable to suppress target genes reliably to analyze unknown phenotypes. Consequently, we explored strategies for direct selection of effective transgenic RNAi lines in Chlamydomonas reinhardtii. initially attempted to suppress expression of the Rubisco small subunit multigene family by placing an IR, homologous to the conserved coding sequence, in the 3'UTR of a transgene conferring resistance to bleomycin. However, this approach was fairly inefficient at inducing RNAi as many strains displayed defective transgene integration, resulting in partial or complete deletion of the IR, or low levels of dsRNA expression, presumably due to transcriptional silencing of the integrated IR transgenes. To overcome these problems we designed a system consisting of tandem IR transgenes that consistently triggered co-silencing of a gene with a selectable RNAi-induced phenotype (encoding tryptophan synthase β subunit) and another gene of interest (encoding either Ku80, an RNA-binding protein, or a thioredoxin isoform). We anticipate that this approach will be useful for generating stable hypomorphic epimutants in high-throughput phenotypic screens.

=> d 3 so

L5 ANSWER 3 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3 SO Plant Journal (2004), 40(4), 611-621 CODEN: PLJUED; ISSN: 0960-7412

=> s 13 and epigenet?

L6 26 L3 AND EPIGENET?

=> dup rem 16
PROCESSING COMPLETED FOR L6
L7 18 DUP REM L6 (8 DUPLICATES REMOVED)

=> d 1-10 ti

- L7 ANSWER 1 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The putative RNA-dependent RNA polymerase RDR6 acts synergistically with ASYMMETRIC LEAVES1 and 2 to repress BREVIPEDICELLUS and microRNA 165/166 in Arabidopsis leaf development
- L7 ANSWER 2 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Atypical RNA polymerase subunits required for RNA-directed DNA methylation
- L7 ANSWER 3 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1
- TI BRU1, a novel link between responses to DNA damage and epigenetic gene silencing in Arabidopsis
- L7 ANSWER 4 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Maintenance of CpG methylation is essential for epigenetic inheritance during plant gametogenesis.
- L7 ANSWER 5 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN
- TI RNA-directed DNA methylation in Arabidopsis.
- L7 ANSWER 6 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN DUPLICATE 2
- TI Two regulatory levels of transcriptional gene silencing in Arabidopsis.
- L7 ANSWER 7 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN
- TI HDA6, a putative histone deacetylase needed to enhance DNA methylation induced by double-stranded RNA.
- L7 ANSWER 8 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
- TI ROS1, a repressor of transcriptional gene silencing in Arabidopsis, encodes a DNA glycosylase/lyase
- L7 ANSWER 9 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Homology-dependent gene silencing mechanisms in fungi.
- L7 ANSWER 10 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI A truncated form of the human CAF-1 p150 subunit impairs the maintenance of transcriptional gene silencing in mammalian cells.

=> d 11-18 ti

- L7 ANSWER 11 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI PTGS in plants, a virus resistance mechanism.
 Original Title: L'inactivation epigenetique posttranscriptionnelle chez les vegetaux: Un mecanisme de resistance aux
 virus.

- *L7 ANSWER 12 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 3
- TI Blocking histone deacetylation in Arabidopsis induces pleiotropic effects on plant gene regulation and development
- L7 ANSWER 13 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4
- TI Endogenous targets of transcriptional gene silencing in arabidopsis
- L7 ANSWER 14 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN
- TI Transcriptional gene silencing mutants.
- L7 ANSWER 15 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 5
- TI Disruption of the plant gene MOM releases transcriptional silencing of methylated genes
- L7 ANSWER 16 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- TI Gene silencing: RNA makes RNA makes no protein
- L7 ANSWER 17 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 7
- TI Cytosine methylation at CG and CNG sites is not a prerequisite for the initiation of transcriptional gene silencing in plants, but it is required for its maintenance
- L7 ANSWER 18 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved.

 (2005) on STN
- TI Manipulation of DET1 expression in tomato results in photomorphogenic phenotypes caused by post-transcriptional gene silencing.

=> d 18 ab

- L7 ANSWER 18 OF 18 AGRICOLA Compiled and distributed by the National Agricultural Library of the Department of Agriculture of the United States of America. It contains copyrighted materials. All rights reserved. (2005) on STN
- The tomato HIGH PIGMENT-2 gene encodes an orthologue of the Arabidopsis AB nuclear protein DE-ETIOLATED 1 (DET1). From genetic analyses it has been proposed that DET1 is a negative regulator of light signal transduction, and recent results indicate that it may control light-regulated gene expression at the level of chromatin remodelling. To gain further understanding about the function of DET1 during plant development, we generated a range of overexpression constructs and introduced them into tomato. Unexpectedly, we only observed phenotypes characteristic of DET1 inactivation, i.e. hyper-responsiveness to light. Molecular analysis indicated in all cases that these phenotypes were a result of suppression of endogenous DET1 expression, due to posttranscriptional gene silencing. DET1 silencing was often lethal when it occurred at relatively early stages of plant development, whereas light hyper-responsive phenotypes were obtained when silencing occurred later on. The appearance of phenotypes correlated with the generation of siRNAs but not DNA hypermethylation, and was most efficient when using constructs with mutations in the DET1 coding sequence or with constructs containing only the 3'-terminal portion of the gene. These results indicate an important function for DET1 throughout plant development and demonstrate that silencing of DET1 in fruits results in increased carotenoids, which may have biotechnological potential.

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     of America. It contains copyrighted materials. All rights reserved.
     (2005) on STN
     Plant journal, p. 344-354
     ISSN: 0960-7412
=> s transcriptionally silenced information
            O TRANSCRIPTIONALLY SILENCED INFORMATION
=> s transcriptionally silent information
             4 TRANSCRIPTIONALLY SILENT INFORMATION
=> dup rem 19
PROCESSING COMPLETED FOR L9
             2 DUP REM L9 (2 DUPLICATES REMOVED)
=> d 1-2 ti
L10 ANSWER 1 OF 2 CAPLUS COPYRIGHT 2005 ACS on STN
     Cloning of transcriptionally silenced plant genes
    ANSWER 2 OF 2 AGRICOLA Compiled and distributed by the National
     Agricultural Library of the Department of Agriculture of the United States
     of America. It contains copyrighted materials. All rights reserved.
     (2005) on STN
                                                       DUPLICATE 1
     Endogenous targets of transcriptional gene silencing in Arabidopsis.
=> s ((steimer a?) or (steimer, a?))/au
           10 ((STEIMER A?) OR (STEIMER, A?))/AU
=> dup rem 111
PROCESSING COMPLETED FOR L11
           5 DUP REM L11 (5 DUPLICATES REMOVED)
=> d 1-5 ti
L12 ANSWER 1 OF 5 CAPLUS COPYRIGHT 2005 ACS on STN
     Cell Culture Models of the Respiratory Tract Relevant to Pulmonary Drug
L12
     ANSWER 2 OF 5 AGRICOLA Compiled and distributed by the National
     Agricultural Library of the Department of Agriculture of the United States
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                                                        DUPLICATE 1
     (2005) on STN
     Epigenetic control of plant development: new layers of complexity.
     ANSWER 3 OF 5 CAPLUS COPYRIGHT 2005 ACS on STN
L12
     Cloning of transcriptionally silenced plant genes
     ANSWER 4 OF 5 AGRICOLA Compiled and distributed by the National
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     Agricultural Library of the Department of Agriculture of the United States
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     (2005) on STN
                                                        DUPLICATE 2
     Apomixis in agriculture: the quest for clonal seeds.
L12 ANSWER 5 OF 5 AGRICOLA Compiled and distributed by the National
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     of America. It contains copyrighted materials. All rights reserved.
     (2005) on STN
                                                       DUPLICATE 3
     Endogenous targets of transcriptional gene silencing in Arabidopsis.
=> s ((sheid o?) or (sheid, o?))/au
             0 ((SHEID O?) OR (SHEID, O?))/AU
=> s ((sheid o?) or (sheid, o?))/au
             O ((SHEID O?) OR (SHEID, O?))/AU
```

SO

L9

T.10

TI

L10

TI

ΤI

ΤI

ΤI

TI

L13

L14

```
=> s ((sheid, o?) or (sheid o?))/au
             0 ((SHEID, O?) OR (SHEID O?))/AU
=> s ((paszkowski j?) or (paszkowski, j?))/au
           259 ((PASZKOWSKI J?) OR (PASZKOWSKI, J?))/AU
L16
=> s l16 and silenc?
L17
           53 L16 AND SILENC?
=> s l18 and plant?
L18 NOT FOUND
The L-number entered could not be found. To see the definition
of L-numbers, enter DISPLAY HISTORY at an arrow prompt (=>).
=> s l17 and plant?
           51 L17 AND PLANT?
L18
=> s 118 and transcript?
           33 L18 AND TRANSCRIPT?
L19
=> dup rem 119
PROCESSING COMPLETED FOR L19
            18 DUP REM L19 (15 DUPLICATES REMOVED)
T-20
=> d 1-10 ti
L20 ANSWER 1 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 1
     BRU1, a novel link between responses to DNA damage and epigenetic gene
     silencing in Arabidopsis
L20 ANSWER 2 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
     DNA and histone methylation in plants
ΤI
L20 ANSWER 3 OF 18 AGRICOLA Compiled and distributed by the National
     Agricultural Library of the Department of Agriculture of the United States
     of America. It contains copyrighted materials. All rights reserved.
     (2005) on STN
                                                        DUPLICATE 2
     Two means of transcriptional reactivation within
TT
     heterochromatin.
    ANSWER 4 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
     Formation of stable epialleles and their paramutation-like interaction in
TΙ
     tetraploid Arabidopsis thaliana
    ANSWER 5 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
     Maintenance of CpG methylation is essential for epigenetic inheritance
     during plant gametogenesis.
    ANSWER 6 OF 18 AGRICOLA Compiled and distributed by the National
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     of America. It contains copyrighted materials. All rights reserved.
     (2005) on STN
                                                        DUPLICATE 3
TI
     Two regulatory levels of transcriptional gene silencing
     in Arabidopsis.
    ANSWER 7 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 4
L20
     Depletion of MOM1 in non-dividing cells of Arabidopsis plants
ΤI
     releases transcriptional gene silencing
    ANSWER 8 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
L20
TI
     Cloning of transcriptionally silenced plant
    ANSWER 9 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
ΤI
     Gene involved in epigenetic gene silencing
```

ANSWER 10 OF 18 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on

L20

STN

'TI' Epigenetic developmental mechanisms in plants: Molecules and targets of plant epigenetic regulation.

=> d 11-18 ti

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 (2005) on STN DUPLICATE 5
- TI Gene silencing and DNA methylation processes.
- L20 ANSWER 12 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 6
- TI Endogenous targets of transcriptional gene silencing in arabidopsis
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 (2005) on STN DUPLICATE 7
- TI Transcriptional gene silencing mutants.
- L20 ANSWER 14 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN DUPLICATE 8
- TI Disruption of the plant gene MOM releases transcriptional silencing of methylated genes
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 (2005) on STN
- TI Disruption of the plant gene MOM releases transcriptional silencing of methylated genes.
- L20 ANSWER 16 OF 18 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Release of epigenetic gene **silencing** by trans-acting mutations in Arabidopsis
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 (2005) on STN DUPLICATE 9
- TI Cytosine methylation at CG and CNG sites is not a prerequisite for the initiation of transcriptional gene silencing in plants but it is required for its maintenance.
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 (2005) on STN DUPLICATE 10
- TI Methylation of cytosines in nonconventional methylation acceptor sites can contribute to reduced gene expression.